Random Tessellation Forests: Appendices

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Appendix A: Theorems and proofs

Lemma 1. With the notation established in Section 2.1 of the main text, let $\Lambda \circ \varphi$ be a product measure $\lambda^d \times \lambda_+$ on $S^{d-1} \times \mathbb{R}_{\geq 0}$. If λ^d is symmetric and if λ_+ is the Lebesgue measure, then Λ is a translation invariant measure on the set H of hyperplanes in \mathbb{R}^d .

Proof: We will first show that Λ is also symmetric (*i.e.*, invariant under reflection through the origin). Suppose that $A \subseteq H$ is a Λ -measurable set of hyperplanes.

$$\Lambda(A) = (\lambda^d \times \lambda_+)(\varphi^{-1}A) \tag{1}$$

$$= \int_0^\infty \lambda^d \left(\{ \overrightarrow{n} : (\overrightarrow{n}, u) \in \varphi^{-1} A \} \right) du \tag{2}$$

$$= \int_0^\infty \lambda^d \left(\left\{ -\vec{n} : (\vec{n}, u) \in \varphi^{-1} A \right\} \right) du = \Lambda(-A). \tag{3}$$

We use the symmetry of λ^d in (3), and in (2) we use the definition of product measures [1].

Let $x \in \mathbb{R}^d$. If $h \in A$, and $\varphi^{-1}h = (\overrightarrow{n}, u)$, then $h = \{P : \langle \overrightarrow{n}, P - u \, \overrightarrow{n} \rangle = 0\}$ and $h + x = \{P : \langle \overrightarrow{n}, P - (u + \langle x, \overrightarrow{n} \rangle) \, \overrightarrow{n} \rangle = 0\}$. Let $A^+ + x = \{h + x \in A + x : \varphi^{-1}h = (\overrightarrow{n}, u) \text{ and } u + \langle x, \overrightarrow{n} \rangle \geq 0\}$ and let $A^- + x = \{h + x \in A + x : \varphi^{-1}h = (\overrightarrow{n}, u) \text{ and } u + \langle x, \overrightarrow{n} \rangle < 0\}$. By these definitions, $(A^+ + x) \cup (A^- + x) = A + x$.

$$\Lambda(A+x) = \Lambda(A^{+}+x) + \Lambda(A^{-}+x) = \Lambda(A^{+}+x) + \Lambda(-(A^{-}+x))$$
(4)

$$= \int_{\overrightarrow{n} \in S^{d-1}} \lambda_{+}(\{u : (\overrightarrow{n}, u) \in \varphi^{-1}(A^{+} + x)\}) + \lambda_{+}(\{u : (\overrightarrow{n}, u) \in \varphi^{-1} - (A^{-} + x)\}) d\lambda^{d}(\overrightarrow{n})$$
 (5)

$$= \int_{\overrightarrow{n} \in S^{d-1}} \lambda_{+}(\{u + \langle x, \overrightarrow{n} \rangle : (\overrightarrow{n}, u) \in \varphi^{-1}A^{+}\}) + \lambda_{+}(\{u + \langle x, \overrightarrow{n} \rangle : (\overrightarrow{n}, u) \in \varphi^{-1} - A^{-})\}) d\lambda^{d}(\overrightarrow{n})$$

$$\tag{6}$$

 $= \int_{\overrightarrow{n} \in S^{d-1}} \lambda_{+}(\{u : (\overrightarrow{n}, u) \in A^{+}\}) + \lambda_{+}(\{u : (\overrightarrow{n}, u) \in -A^{-}\}) d\lambda^{d}(\overrightarrow{n})$ (7)

$$= \Lambda(A^{+}) + \Lambda(-A^{-}) = \Lambda(A^{+}) + \Lambda(A^{-}) = \Lambda(A). \tag{8}$$

Thus, Λ is translation invariant. In (4) and (8), we use the symmetry of Λ (from the first paragraph of this proof). In (7), we use the translation invariance of λ_+ and in (5), we use the definition of product measures [1]. In (6), we use the definitions of $A^+ + x$ and $A^- + x$. Note that $(\overrightarrow{n}, u) \in A^+ + x \Leftrightarrow (\overrightarrow{n}, u + \langle x, \overrightarrow{n} \rangle) \in A^+$ and $(\overrightarrow{n}, u) \in -(A^- + x) \Leftrightarrow (\overrightarrow{n}, u + \langle x, \overrightarrow{n} \rangle) \in -A^-$.

The intuition behind this proof is that translation by any x in A induces a measure preserving shear in $\Lambda \circ \varphi$, as in the stochastic geometry chapter of [2] (this is shown for d=2 in that chapter). In addition, the problem that arises from the fact that translation of a hyperplane through the origin in H flips the sign of \overline{n} is handled by lifting the symmetry of λ^d in S^{d-1} to symmetry of Λ , which requires the decomposition of A+x into a subsets with and without the sign flip (the subsets A^++x , and A^-+x resp.).

Lemma 2. The likelihood function of the labels Z given the tessellation \triangle_t and the labels V and the hyperparameter α can be computed as follows:

$$P(\boldsymbol{Z}|\diamondsuit_{t}, \boldsymbol{V}, \boldsymbol{\alpha}) = \int \cdots \int P(\boldsymbol{Z}, \{\phi_{j}\}_{1 \leq j \leq J_{t}}|\diamondsuit_{t}, \boldsymbol{\alpha}) \, d\phi_{1} \cdots d\phi_{J_{t}}$$
(9)

$$= \int \cdots \int \prod_{i=1}^{J_t} P(\boldsymbol{\phi}_j) \prod_{i=1}^n P(z_i| \mathcal{D}_t, \boldsymbol{\phi}_{h(\boldsymbol{v}_i)}) d\boldsymbol{\phi}_1 \cdots d\boldsymbol{\phi}_{J_t}$$
(10)

$$= \int \cdots \int \prod_{j=1}^{J_t} P(\boldsymbol{\phi}_j) \prod_{i:h(\boldsymbol{v}_i)=j} \prod_{k=1}^K \phi_{jk}^{\delta(z_i=k)} d\boldsymbol{\phi}_1 \cdots d\boldsymbol{\phi}_{J_t}. \tag{11}$$

In (9), we use the conditional independence between Z and \diamondsuit_t . Let $m_{jk} = \sum_{i:h(v_i)=j} \delta(z_i = k)$, and let $m_j = (m_{jk})_{1 \le k \le K}$.

$$P(\boldsymbol{Z}|\boldsymbol{\mathcal{D}}_t) = \int \cdots \int \prod_{j=1}^{J_t} P(\boldsymbol{\phi}_j) \prod_{k=1}^K \phi_{jk}^{m_{jk}} d\boldsymbol{\phi}_1 \cdots d\boldsymbol{\phi}_{J_t}$$
(12)

$$= \int \cdots \int \prod_{i=1}^{J_t} \frac{1}{B(\alpha)} \prod_{k=1}^K \phi_{jk}^{\alpha_k - 1} \prod_{k=1}^K \phi_{jk}^{m_{jk}} d\phi_1 \cdots d\phi_{J_t}$$
 (13)

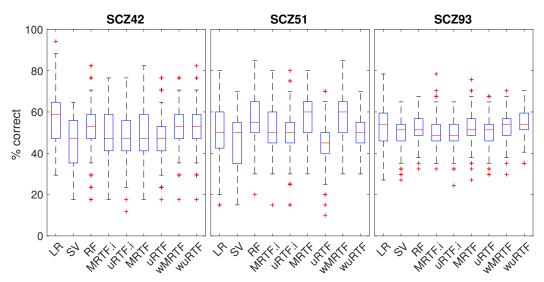
$$=\prod_{j=1}^{J_t} \frac{B(\boldsymbol{\alpha} + \boldsymbol{m}_j)}{B(\boldsymbol{\alpha})}.$$
 (14)

Here $B(\cdot)$ is the multivariate beta function [3].

Lemma 3. In the notation from Section 2.1 of the main text, if Λ is the uRTP measure such that $\Lambda \circ \varphi = \lambda^d \times \lambda_+$ and λ^d is measure associated with the uniform distribution on λ^d , and if B(x,r) is the closed ball centred at x with radius r, then $\Lambda([B(x,r)]) = r$.

Proof: By Lemma 1, the measure Λ is translation invariant, and so $\Lambda([B(x,r)]) = \Lambda([B(0,r)])$. By the definition of product measures [1], $\varphi^{-1}[B(0,r)] = S^{d-1} \times [0,r]$, and so $\Lambda([B(0,r)])$ can be evaluated: $\Lambda([B(0,r)]) = \lambda^d(S^{d-1}) \cdot \lambda_+([0,r]) = r$.

Appendix B: Supplementary figures and tables



Supplementary Figure 1: Boxplots for percent correct on all methods considered for the SCZ42, SCZ51 and SCZ93 datasets. Variance and disorganisation of these results may be due to noise, disorder and difficulty in the mapping from gene expression to schizophrenia. Despite this variance and disorganisation, many of the boxplots show significant improvements over the baseline. For example, a conservative sign test for the improvement of the wuRTF over the baseline in SCZ93 is significant with a p-value of 5.1×10^{-10} .

GL85	BL	LR	SVM	RF	MRTF.i	uRTF.i	MRTF	uRTF	wMRTF	wuRTF
BL		6.7e-18	1.0e+00	7.0e-04	1.0e+00	1.0e+00	1.7e-24	1.0e+00	2.2e-28	5.3e-50
LR	6.7e-18		6.7e-18	1.7e-24	2.3e-20	2.6e-17	2.6e-43	9.9e-20	5.3e-50	8.3e-55
SVM	1.0e+00	6.7e-18		7.0e-04	1.0e+00	1.0e+00	1.7e-24	1.0e+00	2.2e-28	5.3e-50
RF								2.0e-02	1.4e-14	1.2e-37
MRTF.i	1.0e+00	2.3e-20	1.0e+00	1.1e-01		1.0e+00	2.3e-20	1.0e+00		1.5e-44
uRTF.i								1.0e+00	4.6e-30	5.3e-50
MRTF	1		1.7e-24					2.3e-22		2.3e-20
uRTF			1.0e+00						1.4e-27	1.5e-48
wMRTF	1		2.2e-28							3.2e-09
wuRTF	5.3e-50	8.3e-55	5.3e-50	1.2e-37	1.5e-44	5.3e-50	2.3e-20	1.5e-48	3.2e-09	
SCZ42	BL	LR	SVM	RF	MRTF.i	uRTF.i	MRTF	uRTF	wMRTF	wuRTF
BL		4.0e-08	1.0e+00	1.1e-01	9.9e-01	1.0e+00	8.2e-01	1.0e+00	1.8e-03	1.3e-05
LR	4.0e-08		4.0e-08	3.8e-02	4.6e-05	4.2e-04	4.4e-03	6.9e-06		3.1e-01
SVM	1.0e+00	4.0e-08		1.1e-01	9.9e-01	1.0e+00	8.2e-01	1.0e+00	1.8e-03	1.3e-05
RF	1.1e-01	3.8e-02	1.1e-01		5.8e-01	2.6e-01	8.9e-01	3.1e-01	9.6e-01	5.3e-01
MRTF.i							1.0e+00	9.6e-01	3.8e-02	5.2e-02
uRTF.i			1.0e+00					1.0e+00		7.0e-04
MRTF			8.2e-01					7.4e-01		1.8e-01
uRTF			1.0e+00						6.9e-02	8.2e-05
wMRTF			1.8e-03							9.5e-01
wuRTF	1.3e-05	3.1e-01	1.3e-05	5.3e-01	5.2e-02	7.0e-04	1.8e-01	8.2e-05	9.5e-01	
SCZ51	BL	LR	SVM	RF	MRTF.i	uRTF.i	MRTF	uRTF	wMRTF	wuRTF
BL		1.1e-01	1.0e+00	7.6e-11	2.8e-03	1.0e+00	1.4e-13	1.0e+00	1.2e-15	4.7e-01
LR	1.1e-01		1.1e-01	2.8e-02	9.3e-01	3.1e-01	1.1e-03	1.4e-04	8.2e-05	2.6e-01
SVM	1.0e+00	1.1e-01		7.6e-11	2.8e-03	1.0e+00	1.4e-13	1.0e+00	4.1e-15	4.7e-01
RF	7.6e-11	2.8e-02	7.6e-11		8.9e-02	4.0e-08	9.8e-01	1.4e-14	9.3e-01	6.9e-06
MRTF.i			2.8e-03				2.8e-03	3.5e-06	8.2e-05	4.2e-01
uRTF.i			1.0e+00					8.2e-01	3.2e-09	9.6e-01
MRTF			1.4e-13					4.2e-19	1.0e+00	2.0e-07
uRTF			1.0e+00						3.4e-16	2.8e-02
wMRTF			4.1e-15							4.4e-14
wuRTF	4.7e-01	2.6e-01	4.7e-01	6.9e-06	4.2e-01	9.6e-01	2.0e-07	2.8e-02	4.4e-14	
SCZ93	BL	LR	SVM	RF	MRTF.i	uRTF.i	MRTF	uRTF	wMRTF	wuRTF
BL		8.9e-02	1.0e+00	3.8e-02	9.6e-01	9.5e-01	6.9e-02	9.7e-01	6.6e-03	5.1e-10
LR	1					0.7- 02	2 6 2 01	0.0.00	0.0	3.6e-01
	8.9e-02		3.1e-01	7.8e-01	1.8e-01	9.7e-03	5.06-01	8.9e-02	8.2e-01	3.06-01
SVM		3.1e-01				9.7e-03 1.0e+00		8.9e-02 1.0e+00		4.0e-08
	1.0e+00			3.6e-01	9.9e-01	1.0e+00	5.8e-01		1.4e-01	
SVM	1.0e+00 3.8e-02	7.8e-01		3.6e-01	9.9e-01 3.1e-01	1.0e+00	5.8e-01 9.1e-01	1.0e+00	1.4e-01 9.3e-01	4.0e-08
SVM RF	1.0e+00 3.8e-02 9.6e-01	7.8e-01 1.8e-01	3.6e-01	3.6e-01 3.1e-01	9.9e-01 3.1e-01	1.0e+00 3.1e-01 9.1e-01	5.8e-01 9.1e-01 3.6e-01	1.0e+00 1.4e-01	1.4e-01 9.3e-01 1.8e-03	4.0e-08 6.9e-02
SVM RF MRTF.i	1.0e+00 3.8e-02 9.6e-01 9.5e-01	7.8e-01 1.8e-01 9.7e-03	3.6e-01 9.9e-01	3.6e-01 3.1e-01 3.1e-01	9.9e-01 3.1e-01 9.1e-01	1.0e+00 3.1e-01 9.1e-01	5.8e-01 9.1e-01 3.6e-01	1.0e+00 1.4e-01 7.8e-01	1.4e-01 9.3e-01 1.8e-03 3.8e-02	4.0e-08 6.9e-02 2.0e-07
SVM RF MRTF.i uRTF.i	1.0e+00 3.8e-02 9.6e-01 9.5e-01 6.9e-02	7.8e-01 1.8e-01 9.7e-03 3.6e-01	3.6e-01 9.9e-01 1.0e+00	3.6e-01 3.1e-01 3.1e-01 9.1e-01	9.9e-01 3.1e-01 9.1e-01 3.6e-01	1.0e+00 3.1e-01 9.1e-01 2.6e-01	5.8e-01 9.1e-01 3.6e-01 2.6e-01	1.0e+00 1.4e-01 7.8e-01 8.6e-01 2.2e-01	1.4e-01 9.3e-01 1.8e-03 3.8e-02	4.0e-08 6.9e-02 2.0e-07 8.2e-05
SVM RF MRTF.i uRTF.i MRTF	1.0e+00 3.8e-02 9.6e-01 9.5e-01 6.9e-02 9.7e-01	7.8e-01 1.8e-01 9.7e-03 3.6e-01 8.9e-02	3.6e-01 9.9e-01 1.0e+00 5.8e-01	3.1e-01 3.1e-01 9.1e-01 1.4e-01	9.9e-01 3.1e-01 9.1e-01 3.6e-01 7.8e-01	1.0e+00 3.1e-01 9.1e-01 2.6e-01 8.6e-01	5.8e-01 9.1e-01 3.6e-01 2.6e-01	1.0e+00 1.4e-01 7.8e-01 8.6e-01 2.2e-01	1.4e-01 9.3e-01 1.8e-03 3.8e-02 6.9e-01	4.0e-08 6.9e-02 2.0e-07 8.2e-05 2.8e-03

Supplementary Table 1: Pairwise sign tests among all methods considered on GL85, SCZ42, SCZ51 and SCZ93 datasets. Each table shows raw sign test p-values indicated by methods in row and column headers (not corrected for multiple testing). Sign tests are conducted in a conservative manner, in which the sign test is one-tailed towards the better method, and ties are assigned in favour of the method that is not better. Bolding in Table 1 of the main text is found by examining the column of this table corresponding to the best method for a dataset, and then bolding every method in that column with p-value > 0.05 (i.e., nominal significance). BL indicates a baseline method in which the mode label in the training dataset is predicted for all data items.

Dataset	BL	LR	SVM	RF	MRTF.i	uRTF.i	MRTF	uRTF	wMRTF	wuRTF
GL85	5.58	10.52	5.58	7.18	6.09	5.16	6.43	5.78	6.16	6.05
SCZ42	10.62	13.89	10.60	12.17	10.93	11.10	11.76	11.40	11.70	11.41
SCZ51	10.65	12.77	10.67	11.74	11.36	11.12	9.84	10.21	11.07	9.69
SCZ93	7.74	9.65	7.24	7.42	7.20	6.94	7.65	7.24	6.86	6.90

Supplementary Table 2: Standard deviation of percent correct on gene expression datasets over 200 random train/test splits across different methods.

Dataset	LR	SVM	RF	MRTF.i	uRTF.i	MRTF	uRTF	wMRTF	wuRTF
GL85	<.08	<.08	0.006	0.003	43.221	21.681	22.614	15.595	14.679
SCZ42	<.08	<.08	0.005	0.001	2.118	5.468	4.675	5.180	5.066
SCZ51	<.08	<.08	0.004	0.002	5.305	8.420	7.764	8.161	8.641
SCZ93	0.006	0.012	0.006	0.005	89.106	43.534	35.802	41.759	41.068

Supplementary Table 3: Comparison of mean running time (in minutes) on gene expression datasets over 200 random train/test splits across different methods. The experiments were run on an Intel Xeon CPU E5-2683v4@2.10GHz.

Appendix C: Algorithm for Poisson thinning based exact inference

In this appendix, we provide a sampling strategy based on Poisson thinning within SMC that is more precise and efficient than the method displayed in Algorithm 2 of the main text. This strategy is listed in Supplementary Algorithm 1.

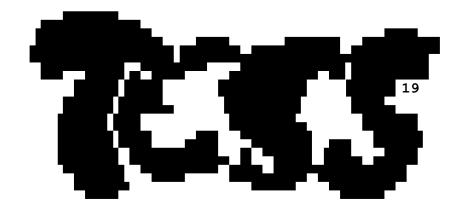
Supplementary Algorithm 1 SMC for inferring RTP posteriors

```
1: Inputs: a) Training dataset V, Z, b) RTP measure \Lambda on H, c) prespecified budget \tau, d) likelihood
      hyperparameter \alpha.
 2: Outputs: Approximate RTP posterior \sum_{m=1}^{M} \varpi_m \delta_{\bigwedge_{\tau}} at time \tau. (\varpi_m are particle weights.)
 3: Set \tau_m \leftarrow 0, for m = 1, \dots, M.
 4: Set \mathcal{O}_{0,m} \leftarrow \{\text{hull } V\}, \varpi_m \leftarrow 1/M, \text{ for } m = 1, \dots, M.
 5: while \min\{\tau_m\}_{m=1}^{M} < \tau do
           Resample \bigwedge_{\tau_m,m}^{m-1} from \{\bigwedge_{\tau_m,m}\}_{m=1}^M w.p.p.t. \{\varpi_m\}_{m=1}^M, for m=1,\ldots,M.
 6:
           7:
           Set \varpi_m \leftarrow 1/M, for m = 1, \ldots, M.
 8:
           for m \in \{m: m=1,\ldots,M \text{ and } \tau_m < \tau\} do
 9:
10:
                Set \tau' \leftarrow \infty, h \leftarrow \emptyset, a \leftarrow \emptyset.
                Set \tau_{a'} \leftarrow \tau_m \ \forall \ a' \in \mathcal{D}_{\tau_m,m}.
11:
                 while 1 do
12:
                      Set \tau^* \leftarrow \infty.
13:
                      for a' \in \mathcal{D}_{\tau_m,m} do
14:
15:
                           if \tau_{a'} < \tau' then
16:
                                 Sample \delta \sim \text{Exp}(r_{a'}).
                                 Sample h' \sim \Lambda(\cdot \cap [B(r_{a'})])/\Lambda([B(r_{a'})]).
17:
                                 Set \tau_{a'} \leftarrow \tau_{a'} + \delta.
18:
19:
                                 if \tau_{a'} < \tau' and h' \cap a' \neq \emptyset then
                                      Set \tau' \leftarrow \tau_{a'}, h \leftarrow h', a \leftarrow a'.
20:
                                 if \tau_{a'} < \tau^* and h' \cap a' = \emptyset then
21:
                                      Set \tau^* \leftarrow \tau_{a'}.
22:
                      if \tau^* > \min\{\tau', \tau\} then
23:
24:
                           break
                25:
26:
                      27:
28:
29:
30:
                      Set \triangle_{t,m} \leftarrow \triangle_{\tau_m,m}, for t \in (\tau_m, \tau].
           Set \tau_m \leftarrow \tau_m + \tau'.
Set \mathcal{Z} \leftarrow \sum_{m=1}^{M} \varpi_m.
31:
32:
33: Set \varpi_m \leftarrow \varpi_m/\mathcal{Z}, for m = 1, ..., M.
34: return the particle approximation \sum_{m=1}^{M} \varpi_m \delta_{\bigwedge_{\tau,m}}.
```

Here, on lines 10 to 24, we provide Poisson thinning for the cutting events. Also, line 12 is an infinite loop, broken by lines 23 and 24 (*i.e.*, it is a **do/until** loop). We note that the computational cost of this method may be reduced by sorting the a' in the loop on line 14 according to the expectation (or a realized value) of the $\tau_{a'}$, or by organising the loops on lines 12 and 14 in a different way in order to respect the cut probabilities. On line 27, h^- and h^+ refer to the two closed half-spaces bounded by h.

Differing from Algorithm 2 in the main text, instead of sampling the time to the next cut, and then choosing the involved polytope w.p.p.t. the cut intensities, we record cuts for all polytopes and find the earliest cut (after thinning). We then discard all future cuts before continuing (making use of the 'add-min' nature of the exponential distribution, as is standard in MJP work). In the thinning code, the variable τ' records the time to the earliest cut, and τ^* records the minimum of the budgets expended by each of the polytopes that have not yet been cut. For more detail on Poisson thinning, we refer to Chapter 2 of [2].

Appendix D: Manual for tess19 v1.0



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The tess19 software implements the Bayesian nonparametric methods described in Ge et al., *Random Tessellation Forests*, 2019. This software constructs a random forest for posterior prediction of categorical data based on real valued predictors. The trees of the random forest are found through SMC inference. This manual is for version v1.0. This software requires the following R packages: optparse, purrr. This software is released under the open source BSD 2-clause license.

1. Basic Usage

```
tess19 <IFILE.txt> <OFILE.txt>
```

Predictions for the missing labels in the Levels column of the file <IFILE.txt> are made using the predictors in the file <IFILE.txt> and the uRTF model, with 100 trees, with the prespecified budget $\tau=\infty$, and with the hyperparameter settings $\alpha_k=n_k/1000$. The predictions are saved in the Levels column of the file <OFILE.txt>.

The format of the file <IFILE.txt> is as follows. The file is space separated. The first line is a header line with one column name for each of predictor (for example, V1 V2 ...), followed by a column named Levels. Subsequent lines are given with one line per data item, with the predictors for the data item followed by the items level. The predictors are real numbers and the levels must be positive integers in the set $1, \ldots K$, where K is the number of levels. Both test and train data must be provided in <IFILE.txt>, and the test data items must have missing labels indicated by the string NaN. The predictors may not have missing data.

The format of the file <OFILE.txt> is as follows. The first line is a header line naming the column Levels (*i.e.*, one column). Subsequent lines are given with one line for data item. If the data item is a training data item, then the value NaN is recorded in the corresponding line. If the data item is a testing data item, then a predicted label is recorded.

2. Advanced Usage

tess19 --usage

tess19 --license

tess19 --version

```
tess19 [--Mondrian] [--weights <WFILE.txt>] [--cuts <MAX-CUTS>]
[--tau <PRESPECIFIED-BUDGET>] [--alpha <HYPER-PARAMETER>] [--trees
<NUMBER-OF-TREES>] [--particles <PARTICLES>] [--seed <SEED>]
<IFILE.txt> <OFILE.txt>
```

- --usage. Prints this manual to the standard output.
- --license. Prints the open source BSD 2-clause license for this software.
- --version. Prints the software version information.
- --Mondrian. Instructs tess19 to conduct axis aligned cuts, yielding the MRTF model, or (if the --weights flag is provided) the wMRTF model. By default, axis aligned cuts are not used.
- --weights <WFILE.txt>. Instructs tess19 to use a weighted version of the uniform distribution for the measure λ^d , yielding the wuRTF or wMRTF model. The weights ω_j are read from the file <WFILE.txt> which must contain d lines corresponding to the prior weight for each predictor, with one real number per line.
- --cuts <MAX-CUTS>. This flag sets a stopping condition wherein SMC particles will return after <MAX-CUTS> cuts, regardless of the budget. By default, <MAX-CUTS> is set to 100. This value must be a positive integer. By the pausing condition, each cut separates a data item, and setting <MAX-CUTS> to a value larger than or equal to the number of data items is equivalent to setting <MAX-CUTS> to ∞ .
- --tau <PRESPECIFIED-BUDGET>. This flag specifies the budget. The budget τ must be a positive real number, or ∞ (the string Inf). By default, τ is set to infinity.
- --alpha <HYPER-PARAMETER>. This flag is a positive real number that sets the coefficient of the empirical label proportion in the Dirichlet/multinomial prior, so that the value of the hyperparameter is $\alpha_k = \text{<HYPER-PARAMETER>} * n_k$. The default value is 10^{-3} .
- --ntrees <NUMBER-OF-TREES>. This flag is a positive integer that sets the number of trees to use in the random forest. By default, <NUMBER-OF-TREES> is set to 100. A value of 1 specifies the uRTP/wuRTP/MRTP/wMRTP priors (i.e., no random forest).
- --particles <PARTICLES>. This flag sets the number of particles to use in the SMC approximations. The default value of <PARTICLES> is 100.
- --seed <SEED>. This flag sets the random seed to <SEED>. The default is to use the system clock to set the random seed.

3. License

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